

Sisson

CRF Errors Corrected by the STIC System Branch

Serial Number: 08/091,240B

CRF Processing Date: 2/26/98
 Edited by: AV
 Verified by: _____ (STIC staff)

ENTERED

#11 1/2

- ☒ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number input by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☒ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: 1-3
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: Seq 2 - moved (ix) FEATURE section

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/791,240BDATE: 02/26/98
TIME: 18:35:24

INPUT SET: S23869.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

1 SEQUENCE LISTING
2
3 (1) General Information
4
5 (i) APPLICANT: Ryncarz, Alexander J.
6
7
8 (ii) TITLE OF THE INVENTION: POSITIVE CONTROLS IN POLYNUCLEOTIDE
9 AMPLIFICATION
10
11
12 (iii) NUMBER OF SEQUENCES: 3
13
14 (iv) CORRESPONDENCE ADDRESS:
15 (A) ADDRESSEE: LINDA M. BUCKLEY
16 DIKE, BRONSTEIN, ROBERTS & CUSHMAN, LLP
17 (B) STREET: P.O. 130 WATER STREET
18 (C) CITY: BOSTON
19 (D) STATE: MA
20 (E) COUNTRY: USA
21 (F) ZIP: 02109-4280
22
23 (v) COMPUTER READABLE FORM:
24 (A) MEDIUM TYPE: Diskette
25 (B) COMPUTER: IBM Compatible
26 (C) OPERATING SYSTEM: DOS
27 (D) SOFTWARE: FastSEQ for Windows Version 2.0
28
29 (vi) CURRENT APPLICATION DATA:
30 (A) APPLICATION NUMBER: 08/791,240
31 (B) FILING DATE: 01/30/1997
32 (C) CLASSIFICATION:
33
34 (vii) PRIOR APPLICATION DATA:
35 (A) APPLICATION NUMBER:
36 (B) FILING DATE:
37
38
39 (viii) ATTORNEY/AGENT INFORMATION:
40 (A) NAME: Buckley, Linda M.
41 (B) REGISTRATION NUMBER: 31,003
42 (C) REFERENCE/DOCKET NUMBER: 47,890 (1008)
43
44 (ix) TELECOMMUNICATION INFORMATION:
45 (A) TELEPHONE: (617) 523-3400

Does Not Comply
Corrected Diskette Needed

ASET
conversion

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/791,240BDATE: 02/26/98
TIME: 18:35:25

INPUT SET: S23869.raw

46 (B) TELEFAX: (617) 523-6440
47
48

ERRORED SEQUENCES FOLLOW:

49 (2) INFORMATION FOR SEQ ID NO:1:
50
51 (i) SEQUENCE CHARACTERISTICS:
52 (A) LENGTH: 25 base pairs
53 (B) TYPE: nucleic acid
54 (C) STRANDEDNESS: single
55 (D) TOPOLOGY: linear
56
57
58 (ii) MOLECULE TYPE: Genomic DNA
59
60 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
61
--> 62 CGACTCACT ATAGGGCGAA TTGGGC
63

(27) 25

64 (2) INFORMATION FOR SEQ ID NO:2:
65
66 (i) SEQUENCE CHARACTERISTICS:
67 (A) LENGTH: 26 base pairs
68 (B) TYPE: nucleic acid
69 (C) STRANDEDNESS: single
70 (D) TOPOLOGY: linear
71
72 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
73
--> 74 CGACTCACT ATAGGGCCGA ATTGGNN
75
76 (i) (ii) FEATURE: N at position 25 is guanine phosphorothioate.
77 N at position 26 is cytosine phosphorothioate.
78
79

(28) 26

80 (2) INFORMATION FOR SEQ ID NO:3:
81
82 (i) SEQUENCE CHARACTERISTICS:
83 (A) LENGTH: 24 base pairs
84 (B) TYPE: nucleic acid
85 (C) STRANDEDNESS: single
86 (D) TOPOLOGY: linear
87
88
89 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
90

PAGE: 3

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/791,240B

DATE: 02/26/98
TIME: 18:35:27

--> 91 CATTAGGCA CCCCAGGCTT TACAC
92

INPUT SET: S23869.raw

26

24

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/791,240BDATE: 02/26/98
TIME: 18:35:28**INPUT SET: S23869.raw**

Line	Error	Original Text
62	# of Sequences for line conflicts w/ running total	CGACTCACT ATAGGGCGAA TTGGGC
74	# of Sequences for line conflicts w/ running total	CGACTCACT ATAGGGCCGA ATTGGNN
91	# of Sequences for line conflicts w/ running total	CATTAGGCA CCCCAGGCTT TACAC